Sage Grouse Landscape Genetics for Management And Conservation

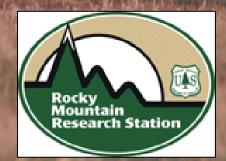
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Sage Grouse Connectivity Outline

- I. Background Connectivity Principles
- II. Sage Grouse Connectivity Project



WHY DO WE CARE ABOUT GENE FLOW AND MOVEMENT OF ANIMALS?



As some of our British parks are ancient, it occurred to me that there must have been long-continued close interbreeding with the fallow-deer (*Cervus dama*) kept in them; but on inquiry I find that it is a common practice to infuse new blood by procuring bucks from other parks.

changed, there can, I think, be no doubt but that a judicious cross with a good stock is of the greatest consequence, and is indeed essential, sooner or later, to the prosperity of every well-ordered park.





LIVESTOCK BREEDERS HAVE KNOWN OF THE DANGERS OF CLOSE RELATED BREEDING

Milk production per cow per lactation increased from **17,444 lbs** to **25,013 lbs** from 1978 to 1998 for the Holstein breed.

High producing cows are increasingly difficult to breed and are subject to **higher health costs** than cows of lower genetic merit for production.



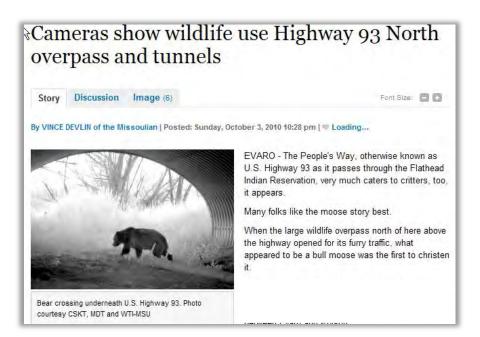
WITHOUT MOVEMENT AND GENE FLOW

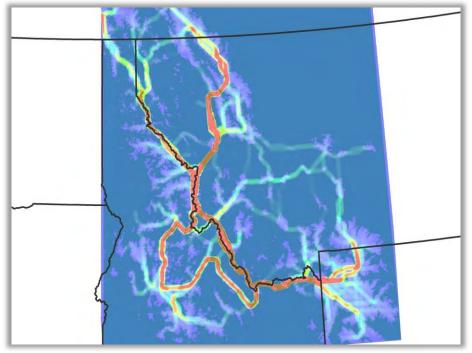
- Inbreeding of Populations
- Loss of genetic diversity
- Depression of Population Fitness
- Suite of Demographic Problems Arise
 (As a consequence of inbreeding or alone)
- Increase in Extinction Risk

SOLUTION: OUTCROSSING – MANY APPROACHES









Most Effective Is through Good Land Overview Stewardship Fostering Connectivity





News is Filled with Actions Aimed at INCREASING WILDLIFE CONNECTIVITY!

Environment ... msnbc.com

More wildlife getting helped across the highway

Funding, studies, new attitu FOXNEWS.COM HOME > U.S.

Protected Path Sought for Wyo. Antelope

Monday, March 03, 2008

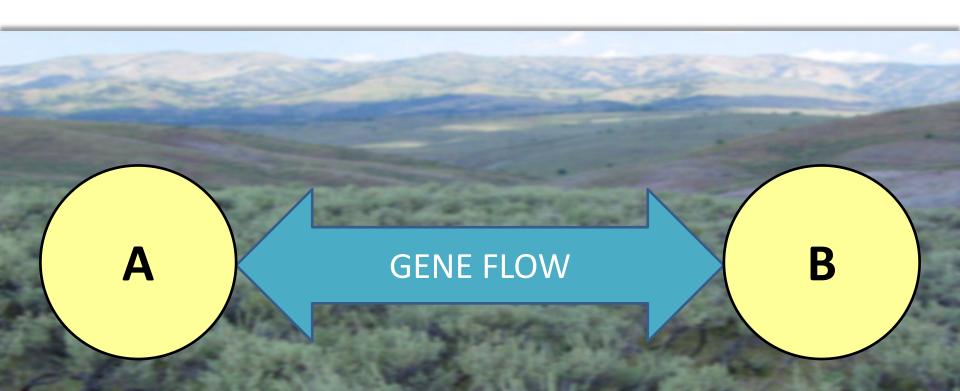
By BOB MOEN, Associated Press Writer





GENE FLOW

GENE FLOW IS THE INCORPORATION OF GENES INTO THE GENE POOL OF ONE POPULATION FROM OTHER POPULATIONS



How Much Gene Flow Is Enough?

The One-Migrant-per-Generation Rule in Conservation and Management

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Abstract: In the face of continuing babitat fragmentation and isolation, the optimal level of connectivity between populations has become a central issue in conservation biology. A common rule of thumb holds that one migrant per generation into a subpopulation is sufficient to minimize the loss of polymorphism and beterozygosity within subpopulations while allowing for divergence in allele frequencies among subpopulations. The one-migrant-per-generation rule is based on numerous simplifying assumptions that may not hold in natural populations. We examine the conceptual and theoretical basis of the rule and consider both genetic and nongenetic factors that influence the desired level of connectivity among subpopulations. We conclude that one migrant per generation is a desirable minimum, but it may be inadequate for many natural populations. We suggest that a minimum of 1 and a maximum of 10 migrants per generation would be an appropriate general rule of thumb for genetic purposes, bearing in mind that factors other than genetics may further influence the ideal level of connectivity.

GENE FLOW

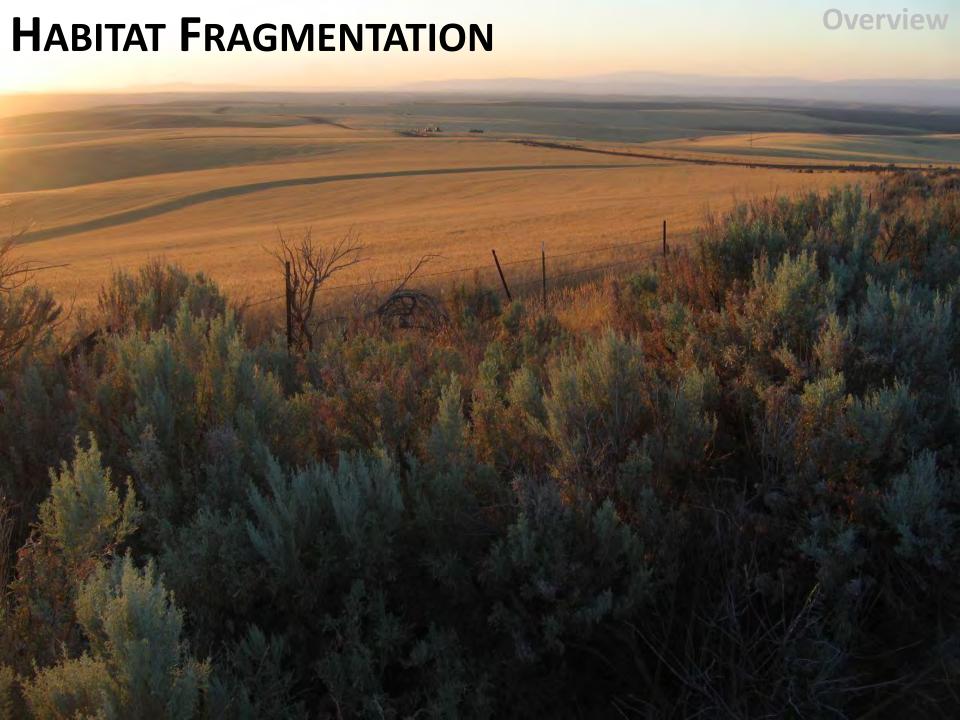
ARE ANIMALS MOVING (AND BREEDING) FROM ONE AREA TO ANOTHER?















HABITAT FRAGMENTATION

Fragmentation & Removal

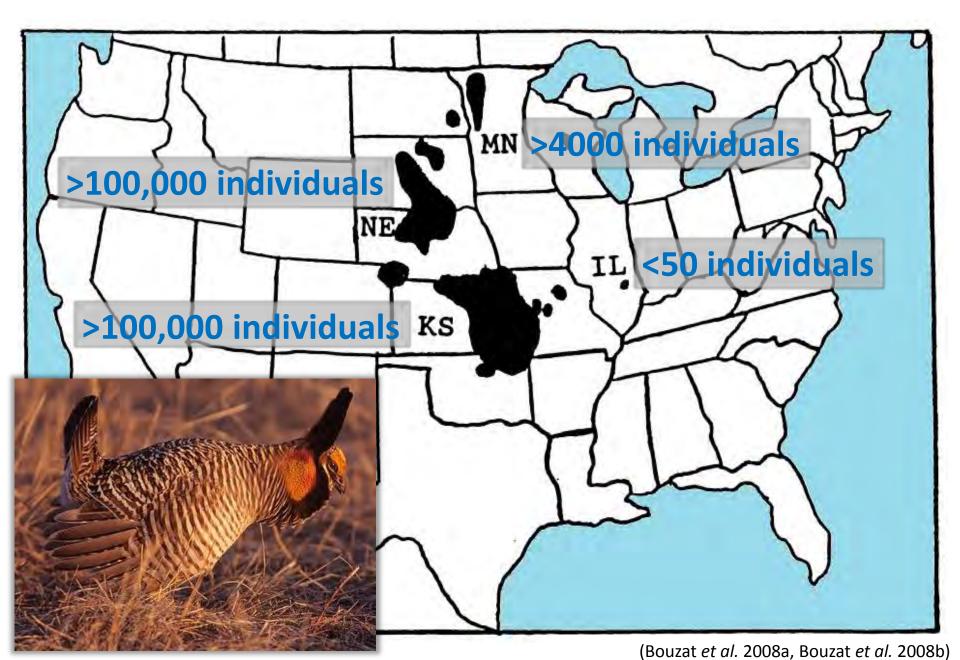
Increased Isolation & Population Size Reduction

Reduced Movement & Inbreeding

Reduced Gene Flow & Loss of Genetic Diversity



GREATER PRAIRIE CHICKEN



GREATER PRAIRIE CHICKEN IN ILLINOIS

Demographic Data

Population Size

1933: 25,000

1962: 2,000

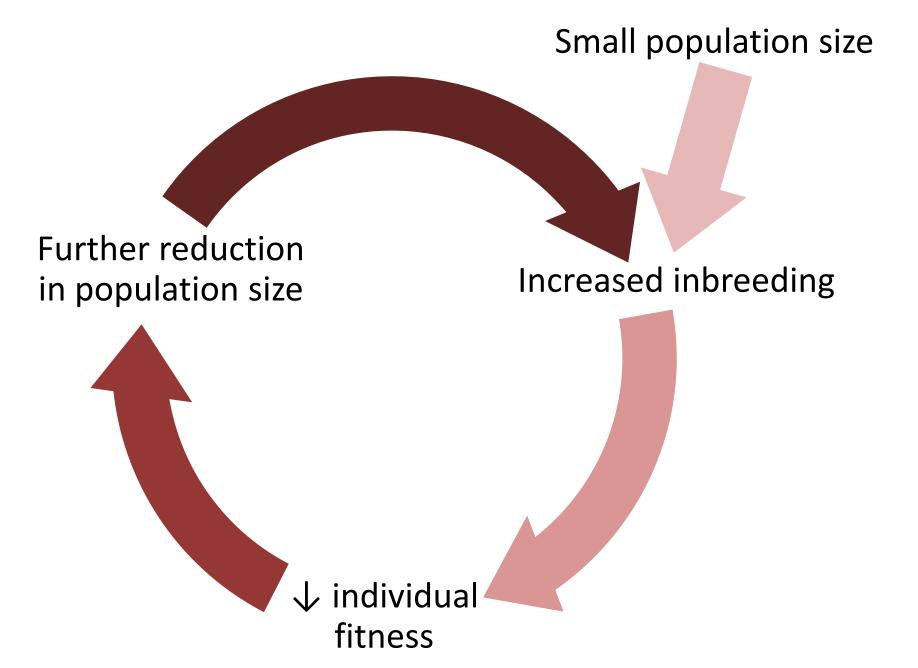
1993: 50

Hatching Success Rate

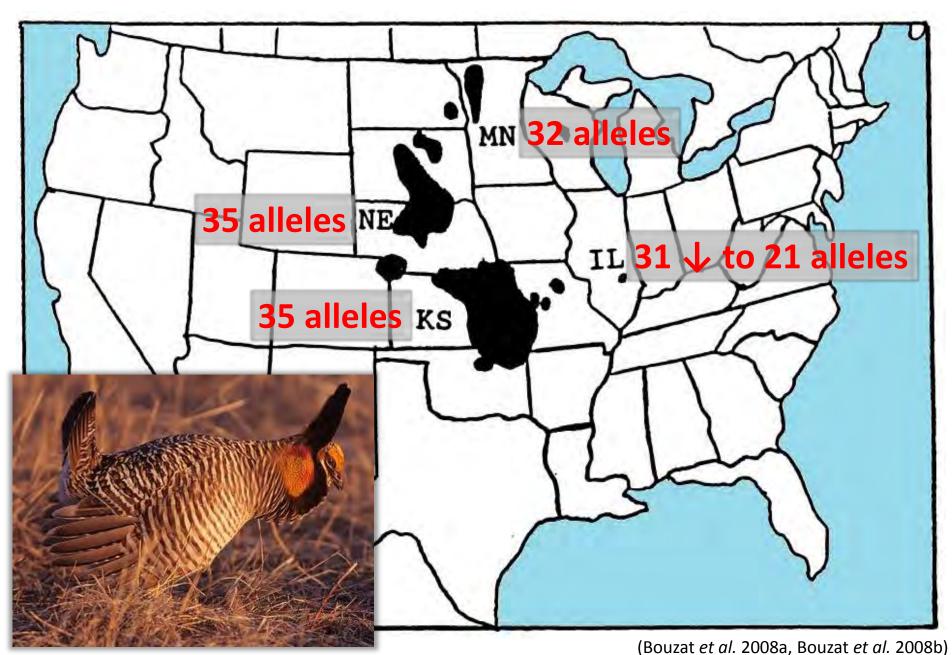
1960 - 90%

1990 - 74%

SMALL POPULATION SIZE



GREATER PRAIRIE CHICKEN IN ILLINOIS



GREATER PRAIRIE CHICKEN IN ILLINOIS

Demographic Data

Population Size

1933: 25,000

1962: 2,000

1993: 50

Hatching Success Rate

1960 - 90%

1990 - 74%

Genetic Data

1960-1990 Genetic Variation Declined 30%

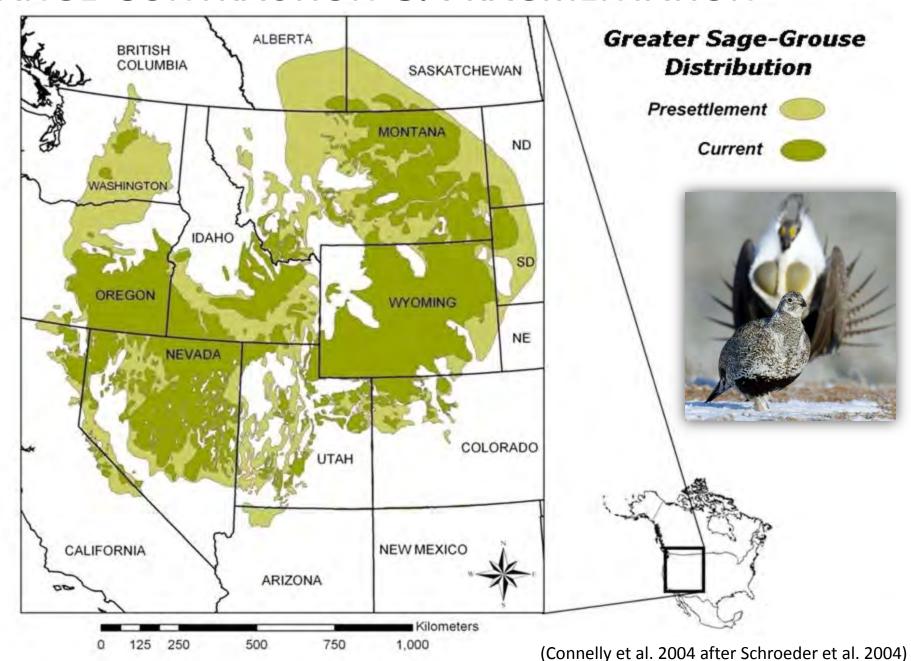
Introduction Effort

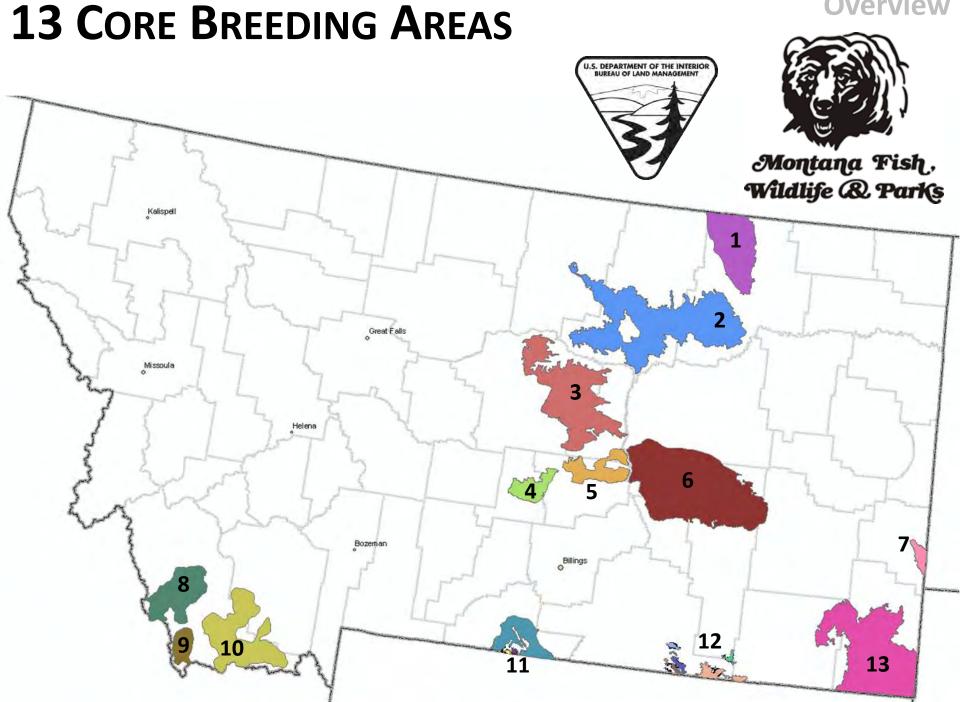
Hatching Rate Success 1 for MN x IL & KA x IL to 94%

(Westemeier et al. 1998)

Overview

RANGE CONTRACTION & FRAGMENTATION

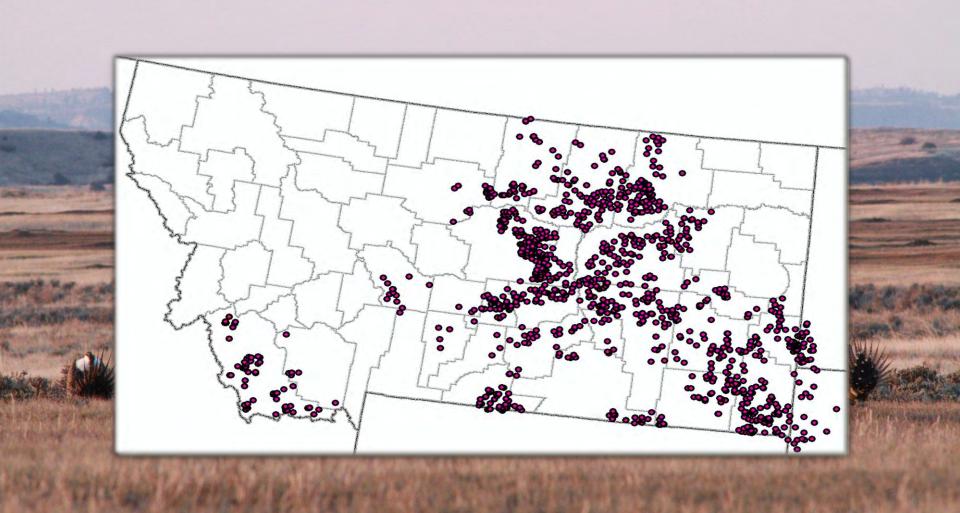


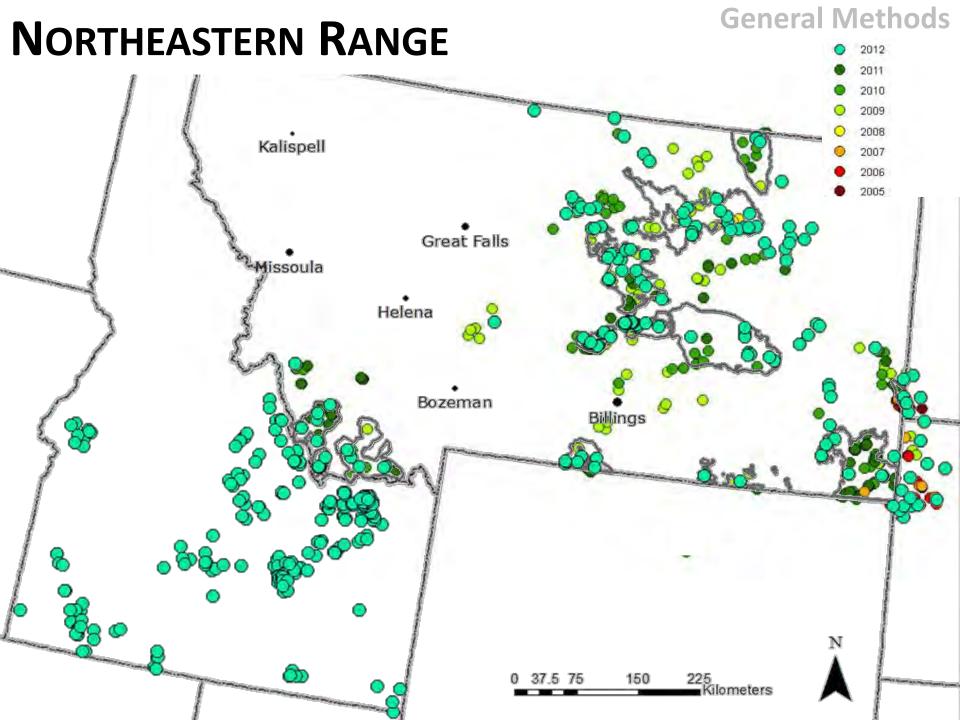


OVERARCHING GOAL:

TO DELINEATE GREATER SAGE-GROUSE
CONSERVATION UNITS TO CONSERVE THE
GENETIC STRUCTURE, CONNECTIVITY, AND
GENETIC VARIATION ACROSS A CHANGING
LANDSCAPE

SAMPLING UNIT - THE LEK





SAMPLING EFFORT

State	Total Leks	Total Feathers	
ID	224	2686	
MT	467	5730	
ND	9	221	
SD	18	318	
TOTAL	718	8955	

RANGE-WIDE GENETICS



















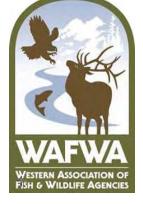








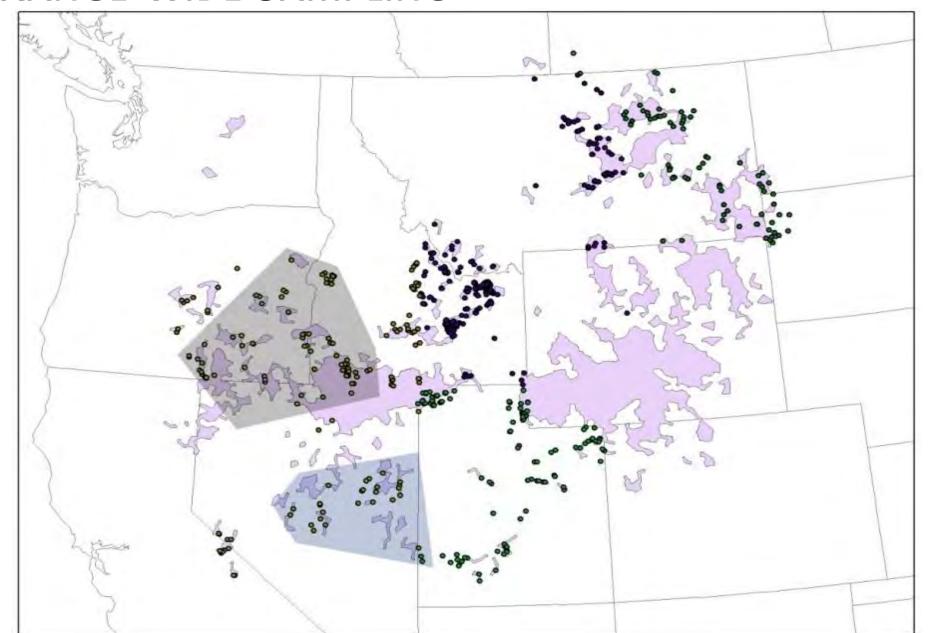








RANGE-WIDE SAMPLING



(image courtesy of Steve Knick)

DNA EXTRACTION & PROCESSING



	Extracted	Amplified	Individuals
Current	3792	2911	1620
Expected	5000	5000	3500

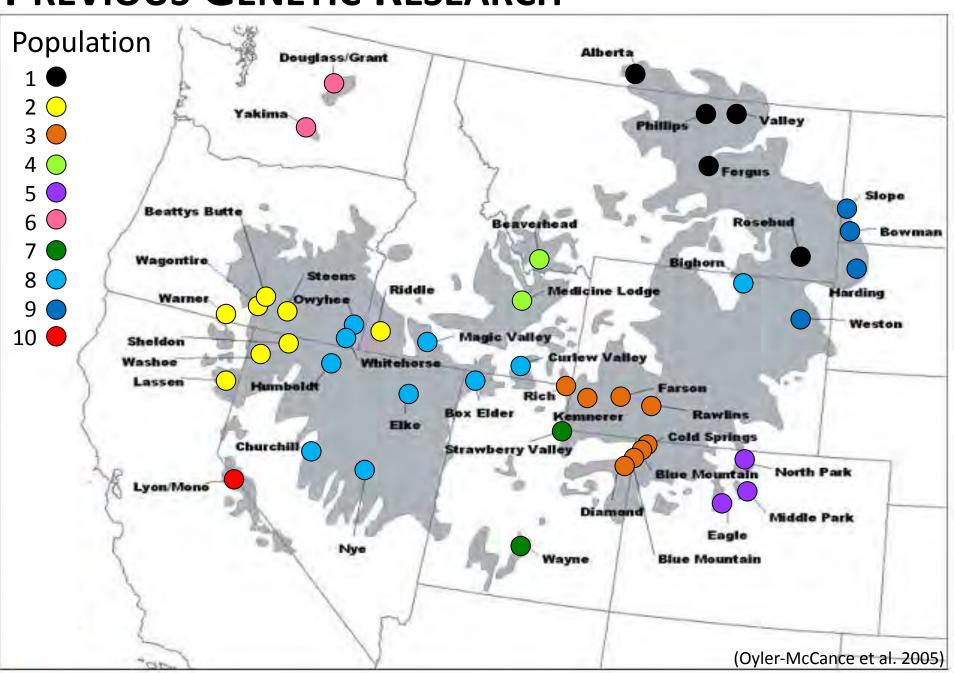
DEVELOP HIGH RESOLUTION GENETIC MARKERS FOR INCREASED POWER OF ANALYSIS.



IDENTIFY POPULATION STRUCTURE.



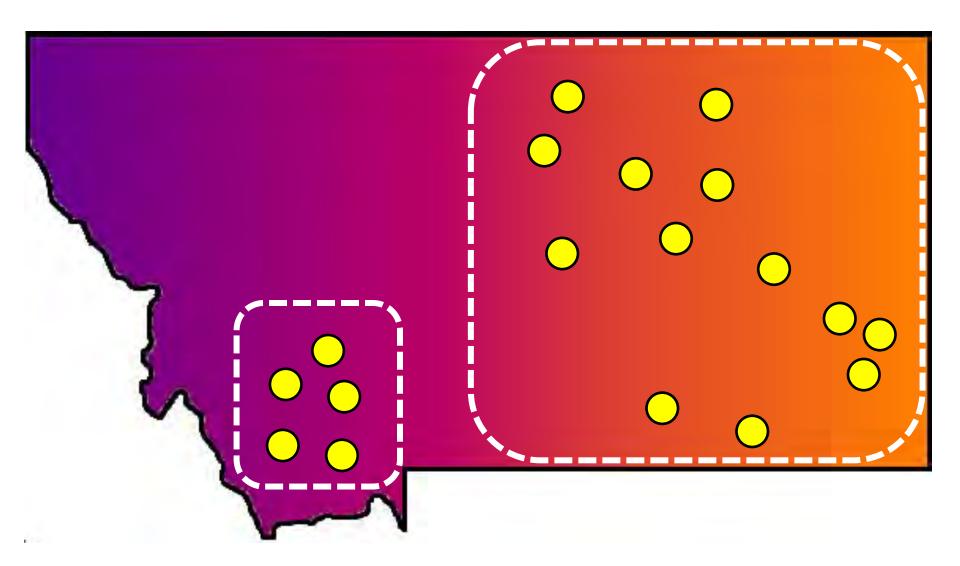
PREVIOUS GENETIC RESEARCH



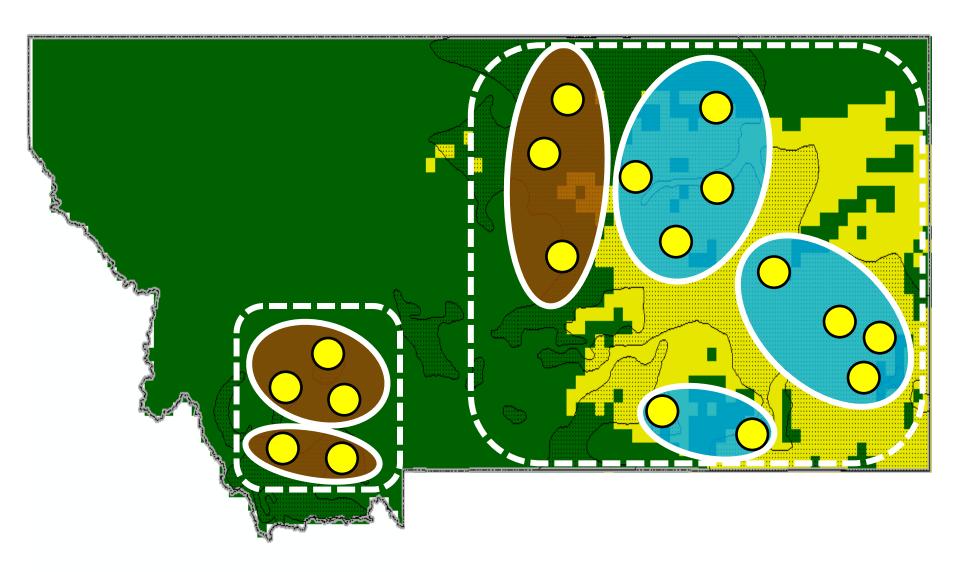
a. Does genetic population structure exist?

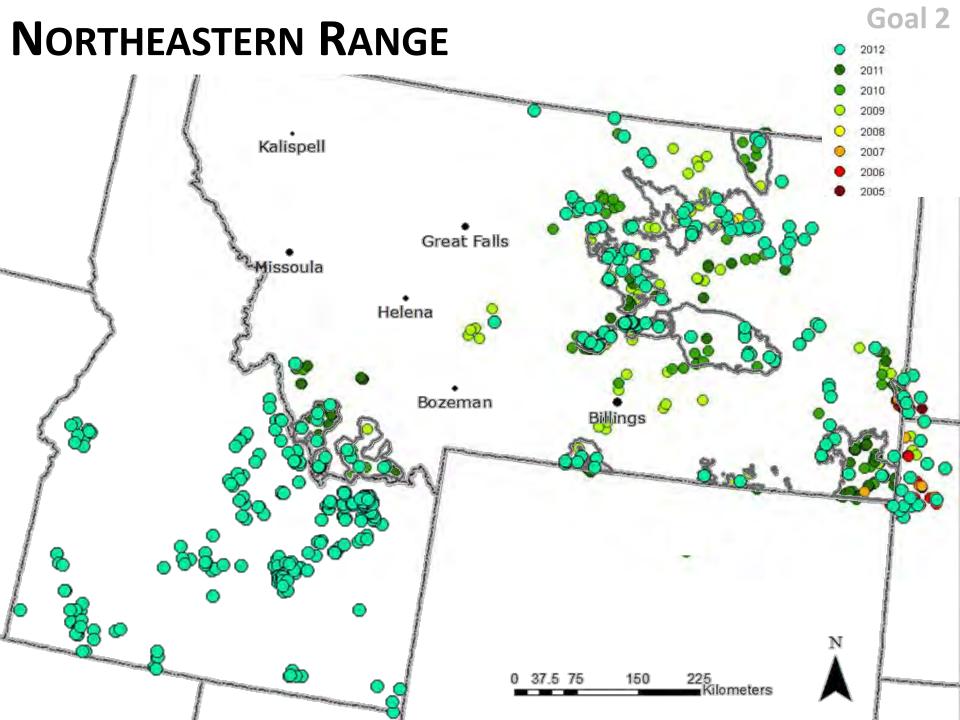


b. Where are the major breaks in genetic continuity?



c. What is the genetic composition of each subpopulation?





ASSESS OF THE EFFECT OF LANDSCAPE FEATURES ON GENE FLOW



LANDSCAPE DISTURBANCE

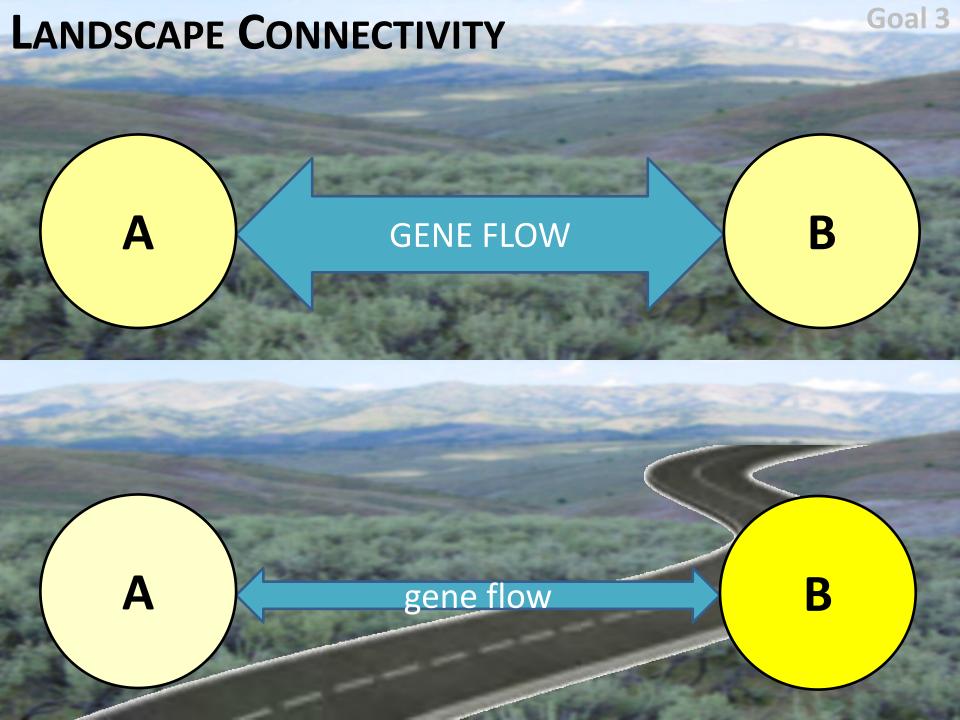
Disturbance reduces lek attendance leads to extirpation

(Knick et al. 2013)

rivers
roads
cultivated cropland
energy development
sagebrush land cover
conifer encroachment
human population density

(Patterson 1952; Remington & Braun 1991; Lyon & Anderson 2003; Holloran & Anderson 2005; Aldridge et al. 2008; Knick & Hanser 2011; Johnson et al. 2011; Johnson et al. 2011)



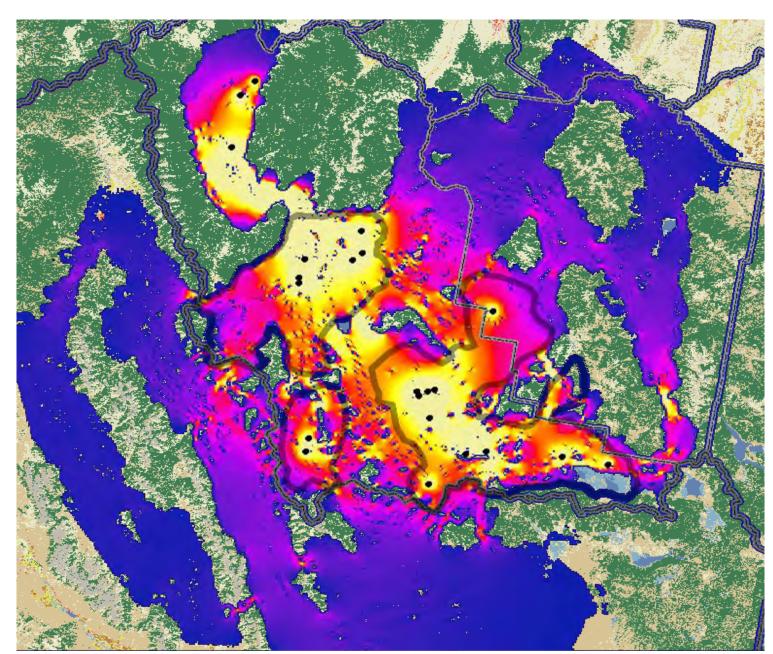


CONNECTIVITY MAP CREATION

Resistance to Movement (Gene Flow)

Low

High

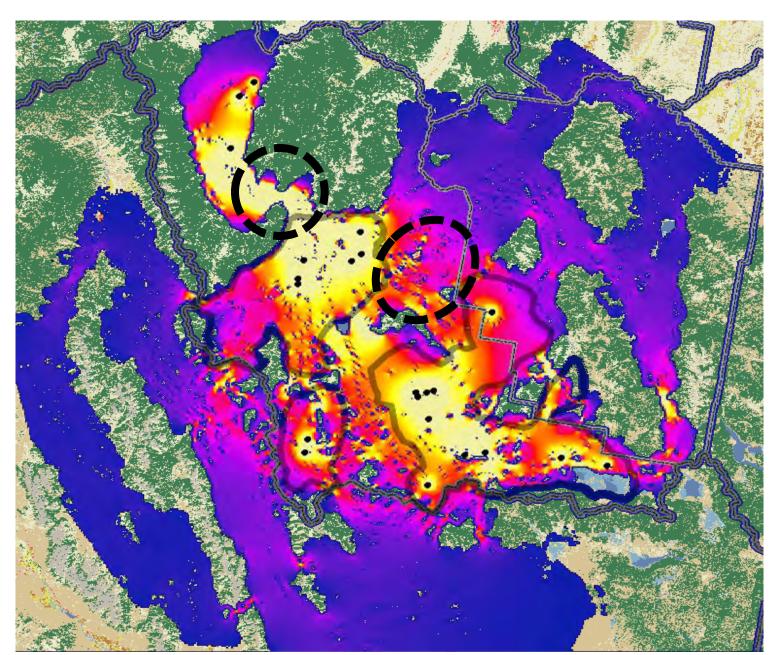


CONNECTIVITY MAP CREATION

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Low

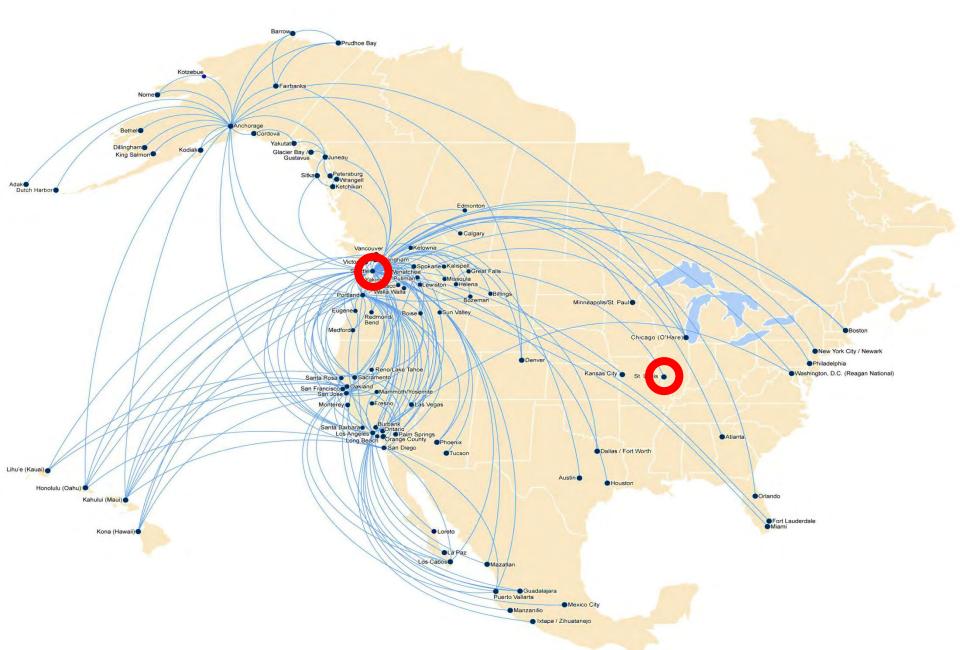
High



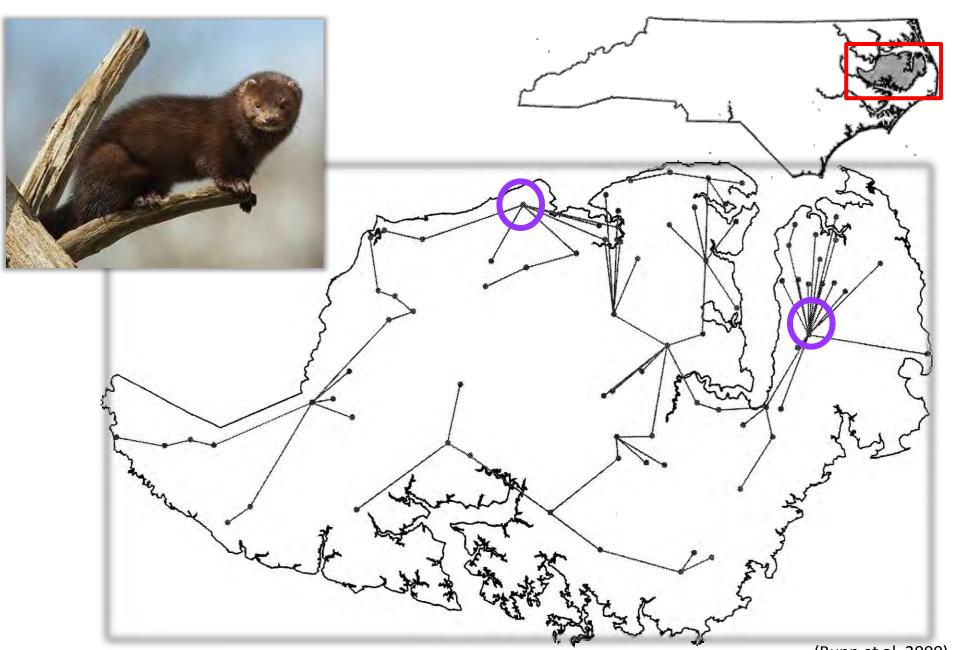
MODEL THE RELATIVE IMPORTANCE OF EACH LEK TO MAINTAINING POPULATION CONNECTIVITY AND PERSISTENCE.



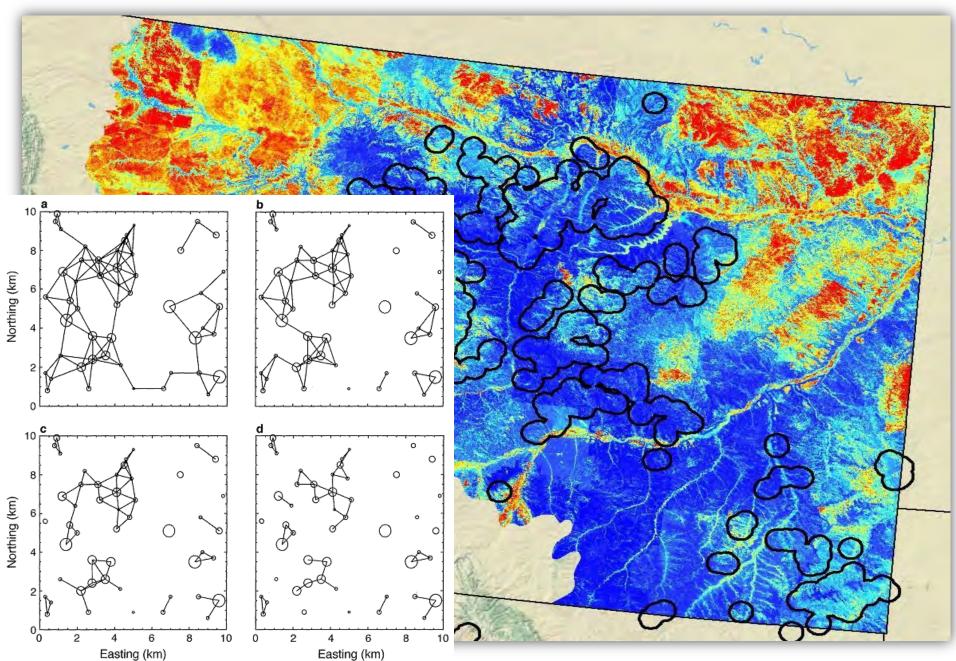
REAL-WORLD NETWORKS



GRAPH STRUCTURE - WILDLIFE



NETWORK RESILIENCY















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